



Sheet 1 of 2

APPLICANT FACSIMILE OF FORM PTO-1449

U.S. DEPARTMENT OF
COMMERCE
PATENT AND TRADEMARK OFFICE

ATTY DOCKET NO

PKZ-043

SERIAL NO.

09/801563

REVISED

LIST OF PUBLICATIONS CITED BY APPLICANT
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APPLICANT

Levy, Stuart B. et al.

FILING DATE

March 8, 2001

GROUP

1648

U.S. PATENT DOCUMENTS

EXAMINER INITIAL		DOCUMENT NUMBER	DATE	NAME	CLASS	SUBCLASS	FILING DATE IF APPROPRIATE
32	A1	5,766,924	06/98	Levy	435	252.3	

FOREIGN PATENT DOCUMENTS

		DOCUMENT NUMBER	DATE	COUNTRY	CLASS	SUBCLASS	TRANSLATION Yes NO

OTHERS (including Author, Title, Date, Pertinent Pages, Etc.)

32	A2		Alekshun, M.N. et al. Regulation of chromosomally mediated multiple antibiotic resistance: the <i>mar</i> regulon. <i>Antimicrob. Agents Chemother.</i> 1997 Oct;41(10):2067-75
32	A3		Aono, R. et al. Involvement of outer membrane protein TolC, a possible member of the <i>mar</i> - <i>sox</i> regulon, in maintenance and improvement of organic solvent tolerance of <i>Escherichia coli</i> K-12. <i>J. Bacteriol.</i> 1998 Feb;180(4):938-44
32	A4		Barbosa, T.M. et al. Genetic analysis of the <i>Shigella flexneri mar</i> locus. 1999. 99 th General Meeting of the American Society for Microbiology (Chicago, IL). Abstract A42, p. 9
32	A5		Barbosa, T.M. et al. Differential expression of over 60 chromosomal genes in <i>Escherichia coli</i> by constitutive expression of MarA. <i>J. Bacteriol.</i> 2000 Jun;182(12):3467-74
32	A6		Blattner, F.R. et al. The complete genome sequence of <i>Escherichia coli</i> K-12. <i>Science.</i> 1997 Sep 5;277(5331):1453-74
32	A7		Cohen, S. P. et al. A multidrug resistance regulatory chromosomal locus is widespread among enteric bacteria. <i>J. Infect. Dis.</i> 1993;168:484-8
32	A8		Cohen, S.P. et al. Genetic and functional analysis of the multiple antibiotic resistance (<i>mar</i>) locus in <i>Escherichia coli</i> . <i>J. Bacteriol.</i> 1993 Mar;175(5):1484-92
32	A9		Gallegos, M.-T. et al. The XylS/AraC family of regulators. <i>Nucleic Acids Res.</i> 1993 Feb 25;21(4):807-10
32	A10		Gambino, L. et al. Overexpression of the MarA positive regulator is sufficient to confer multiple antibiotic resistance in <i>Escherichia coli</i> . <i>J. Bacteriol.</i> 1993 May;175(10):2888-94
32	A11		George, A.M. et al. Gene in the major cotransduction gap of the <i>Escherichia coli</i> K-12 linkage map required for the expression of chromosomal resistance to tetracycline and other antibiotics. <i>J. Bacteriol.</i> 1983 Aug;155(2):541-8
32	A12		Gold, H.S. et al. Antimicrobial-drug resistance. <i>N. Engl. J. Med.</i> 1996 Nov 7;335(19):1445-53
32	A13		Li, H. et al. The periplasmic murein peptide-binding protein MppA is a negative regulator of multiple antibiotic resistance in <i>Escherichia coli</i> . <i>J. Bacteriol.</i> 1999 Aug;181(16):4842-7
32	A14		Martin, R.G. et al. Binding of purified multiple antibiotic-resistance repressor protein (MarR) to <i>mar</i> operator sequences. <i>Proc. Natl. Acad. Sci. U.S.A.</i> 1995 Jun 6;92(12):5456-60
32	A15		Martin, R.G. et al. Autoactivation of the <i>marRAB</i> multiple antibiotic resistance operon by the MarA transcriptional activator in <i>Escherichia coli</i> . <i>J. Bacteriol.</i> 1996 Apr;178(8):2216-23

Examiner

Date Considered

*EXAMINER:

Initial if reference considered, whether or not citation is in conformance with MPEP 609; Draw line through citation if not in conformance and not considered. Include copy of this form with next communication to applicant.



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